

Aneuploid sperm formation in rainbow trout exposed to the environmental estrogen 17 α -ethynylestradiol

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Environmental contaminants that mimic native estrogens (i.e., environmental estrogens) are known to significantly impact a wide range of vertebrate species and have been implicated as a source for increasing human male reproductive deficiencies and diseases. Despite the widespread occurrence of environmental estrogens and recognized detrimental effects on male vertebrate reproduction, no specific mechanism has been determined indicating how reduced fertility and/or fecundity is achieved. Previous studies show that male rainbow trout, *Oncorhynchus mykiss*, exposed to the environmental estrogen 17 α -ethynylestradiol (EE2) before gamete formation and fertilization produce progeny with significantly reduced embryonic survival. To determine whether this observed decrease results from sperm chromosome alterations during spermatogenesis, male rainbow trout were exposed to 10 ng of EE2/l for 50 days. After exposure, semen was collected and sperm aneuploidy levels analyzed with two chromosome markers by fluorescent *in situ* hybridization. *In vitro* fertilizations were also conducted by using control and exposed sperm crossed to eggs from an unexposed female for offspring analysis. Evaluations for nucleolar organizer region number and karyotype were performed on developing embryos to determine whether sperm aneuploidy translated into embryonic aneuploidy. Results conclusively show increased aneuploid sperm formation due to EE2 exposure. Additionally, embryonic cells from propagated progeny of individuals possessing elevated sperm aneuploidy display high levels of embryonic aneuploidy. This study concludes that EE2 exposure in sexually developing male rainbow trout increases levels of aneuploid sperm, providing a mechanism for decreased embryonic survival and ultimately diminished reproductive success in EE2 exposed males.

aneuploidy | EE2 | xenoestrogens

Endocrine disrupting chemicals are found ubiquitously worldwide due to human environmental contamination (1–4). These chemicals comprise several different classes and those with an estrogenic mode of action (i.e., environmental estrogens) are of major concern. In industrialized countries concerns about human health impacts resulting from environmental estrogens have been mounting. These concerns largely resulted from a meta-analysis of sperm counts indicating a nearly 50% reduction between 1940 and 1990 (5). This finding proved controversial with additional analyses supporting (6–11) and contradicting (12–14) the original finding. Additional reports suggest that a rapid increase in testicular germ cell cancers, cryptorchidism, and other congenital anomalies in developed countries may be related to increased environmental estrogen exposure (15–18). Additionally, recent animal model studies have indicated that environmental estrogens are capable of altering epigenetic patterns during early development with resulting gene expression changes, increased cancer susceptibility, and congenital aberrations as well as diminished reproductive capacity, compounding human health concerns (19, 20). These studies, and others, have led to the hypothesis that reduced fertility and/or increased reproductive anomalies result from exposure to environmental estrogens.

Environmental estrogens have a wide variety of chemical structures but are grouped together based on their ability to mimic natural estrogen by interfering with/or binding directly to estrogen receptors (21–24). Through these actions environmental estrogens affect vertebrate reproduction across a wide range of doses causing reduced fertility and fecundity, altered reproductive behavior, gonad morphological changes, and decreased embryonic survival (25–34). Although both sexes are affected, males exhibit the highest degree and number of detrimental effects. Specific male defects caused by environmental estrogen exposures in vertebrates include: intersex (25, 29), diminished sperm count (27, 35–37), genital tract alterations (38), increased germ cell apoptosis (30, 33), and male induced embryonic mortality (31, 39).

Despite the evidence for significant effects of environmental estrogen exposure on male vertebrates, no specific mechanism of action has been determined to explain how reduced fertility and/or fecundity of morphologically and physiologically normal individuals are achieved. Although most rodent model studies show decreased sperm production and/or increased apoptosis of germ cells after exposure (27, 30, 33, 35), others have failed to confirm these findings (40). Additionally, recent *in vitro* exposures of human spermatozoa to catechol estrogens (e.g., quercetin, diethylstilbestrol and pyrocatechol) indicate an impact on sperm DNA integrity through altered redox cycling, but estrogen (17 β -estradiol) and other estrogen analogues (nonylphenol and BPA) do not show this effect (41). Despite this *in vitro* finding using spermatozoa, how these compounds would affect spermatogenesis *in vivo* is unknown.

Fish studies in which male rainbow trout (*Oncorhynchus mykiss*) are exposed *in vivo* to the environmental estrogen 17 α -ethynylestradiol (EE2) show no defects in either testis morphology or sperm motility, but exhibit significantly reduced progeny survival when exposed as late stage juveniles during final sexual maturation (31, 42). In these studies, which specifically evaluated the effect of EE2 exposure on the male germ cell at environmentally relevant concentrations (\approx 10 ng/l) during spermatogenesis (i.e., meiosis), the problem was attributed to qualitative sperm defects. Further evaluations led to the consideration of a possible genetic link affecting embryonic survival. This new hypothesis is based on previous studies in which rainbow trout sperm with fragmented DNA (UV irradiated) were used to fertilize eggs *in vitro*, resulting in an observed increase in embryonic aneuploidy (43, 44). Results from these experiments show a significant reduction in the survival of

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embryos produced with UV irradiated sperm compared with control embryos propagated by using un-irradiated sperm (45). This study indicates that sperm chromosome damage leads to a pattern of embryo death similar to that observed when male parents are exposed to EE2 (31, 42). Additionally, evidence exists that exposure of female mice to low levels of BPA and of maturing oocytes to supraphysiological levels of 2-methoxyestradiol, causes increases in aneuploid oocyte formation (46, 47), although the BPA studies could not be reproduced by another study (40). These observations led us to propose that exposure of sexually maturing male rainbow trout to EE2 results in aneuploid sperm formation, which causes a significant proportion of embryos to die shortly after fertilization. This study used the previously established male rainbow trout experimental model, with fish exposed during the actively meiotic, mid-spermatogenic time point with spermatocytes and spermatids predominant in the gonad before final sexual maturation, to determine: (i) whether an environmentally relevant concentration of EE2 during late spermatogenesis results in the formation of aneuploid sperm and (ii) whether progeny produced from males exposed to EE2 during late spermatogenesis exhibit increased levels of aneuploidy.

Results

Five clonal YY male rainbow trout each were exposed as late stage juveniles immediately before sexual maturation (6,700°d; \approx 1 year 9 months of age) to a nominal concentration of 10 ng of EE2/l for 50 days or to methanol only (solvent control), respectively. The mean measured daily in-flow water concentration was 16.1 ng of EE2/l with a mean GCMS measured EE2 level of 8.9 ng of EE2/l, both falling within 50–100% of nominal values. After exposure semen was collected from all sexually mature individuals and cryopreserved for later analyses (four control and four exposed). The remaining two individuals, one per treatment, did not produce sperm or exhibit secondary sexual characteristics and possessed immature testes upon post-mortem examination. This is not unusual as the majority of male rainbow trout typically attain sexual maturity for the first time at two years of age but a small proportion do not until three years of age.

Fluorescent *in situ* hybridization (FISH) analysis on cryopreserved sperm was performed using two probes, an 18s rDNA probe (Vysis green) hybridizing to chromosome 20 and a 5s rDNA probe (Vysis orange) hybridizing to the Y chromosome (48). Fig. 1 shows examples of normal and aneuploid sperm nuclei evaluated by using FISH for this study. Quantitatively FISH analysis with both probes revealed significant increases in levels of sperm aneuploidy for exposed but not control individuals (Fig. 2). Average combined sperm aneuploidy levels for both probes were 1.2% (control) and 29.1% (exposed), and were significantly different ($P < 0.0001$). Although both Y chromosome and chromosome 20, based on FISH analysis, were equally represented in controls, chromosome 20 aneuploidy was more frequently observed in exposed individuals compared with Y chromosome aneuploidy (Table 1).

In vitro fertilizations were performed by using cryopreserved semen and freshly collected eggs from a single, unexposed female to determine offspring aneuploidy levels. Embryo analysis consisted of nucleolar organizer region (NOR) silver staining and karyotype counts. NOR analysis was performed on 25 control and 30 exposed individuals (Fig. 3). Rainbow trout have a single NOR on chromosome 20 (48, 49). Analysis revealed cell nuclei from the control embryos predominantly exhibiting two NORs per nucleus, consistent with normal diploid rainbow trout. Contrary to this finding, only 73% of the embryos propagated from males exposed to EE2 exhibited two NORs. Of the remaining 27%, 17% expressed one NOR per nucleus, and 10% possessed predominantly three NORs per cell nucleus, consis-

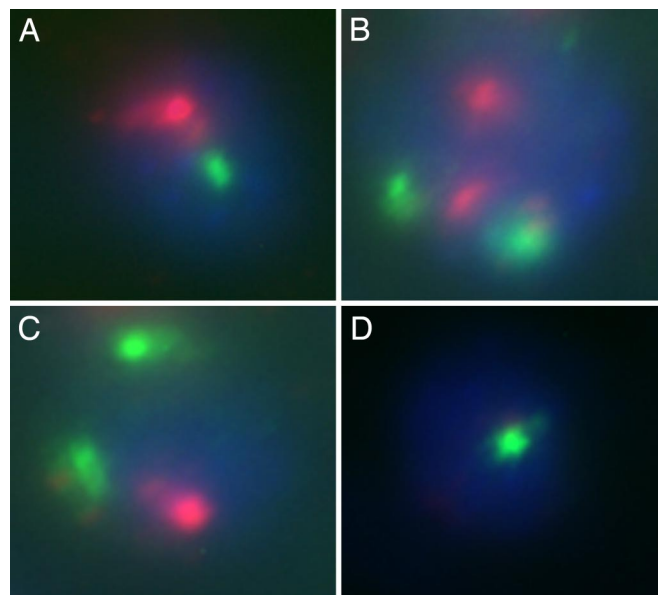


Fig. 1. Representative photomicrographs of normal and aneuploid rainbow trout sperm nuclei with chromosomes identified by using fluorescent *in situ* hybridization. (A) Normal, haploid nucleus (stained blue) from a control sperm with one chromosome 20 (green probe) and one sex chromosome (red probe). (B and C) Examples of hyperploid sperm nuclei and (D) a hypoploid sperm nucleus from EE2 exposed fish.

tent with being haploid (or hypoploid) and triploid (or hyperploid), respectively (Fig. 4A).

Karyotype analysis on 10 randomly selected embryos from the control group revealed a chromosome count of 62 with 104 chromosome arms, which is consistent with previous reports for this population (50). An embryo chromosome count was considered aneuploid if it deviated from controls for either a whole chromosome or a chromosome arm number with a minimum 10 of 15 individual spreads possessing the same count required to determine ploidy (Fig. 5). Analysis of 24 embryos from EE2 exposed males revealed only 42% with normal karyotypes. Among individuals deemed aneuploid 42% were hypoploid and 16% hyperploid. Exposed group embryonic chromosome counts ranged from 57 to 90 with associated arm numbers ranging from 97 to 153, respectively (Fig. 4B). Ten exposed group embryos were analyzed for both NOR and karyotype. Among these ten

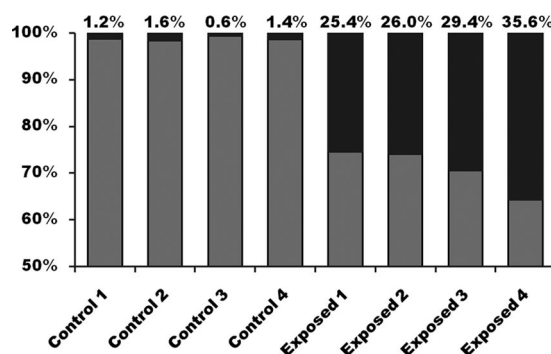


Fig. 2. Combined total percent of normal (gray) and aneuploid (black) sperm collected from control and exposed males evaluated by fluorescent *in situ* hybridization. The percentage of aneuploid sperm for each individual exhibited is given above each bar. Differences between groups were statistically significant at $P < 0.0001$.

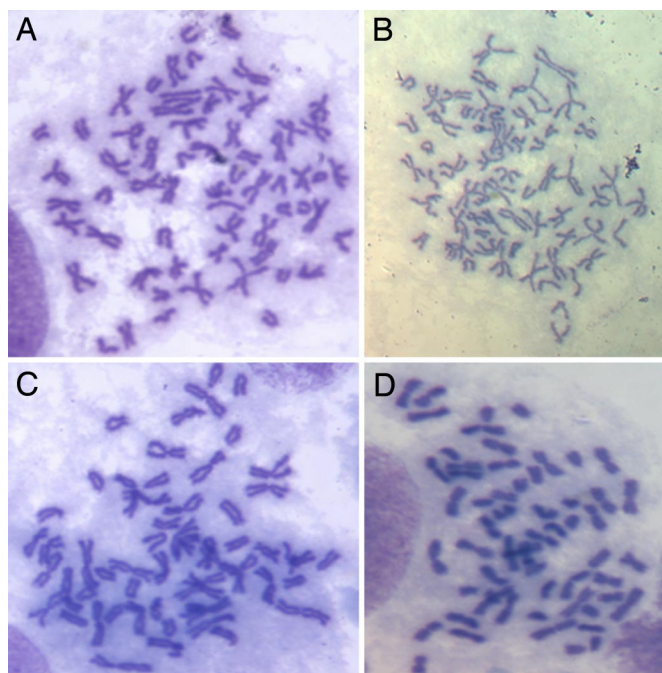


Fig. 5. Photomicrographs showing typical rainbow trout chromosome spreads used for karyotype analysis. (A) Normal, diploid cell; (B) hyperploid cell; and (C and D) hypoploid cells.

recombination sites, or chiasma, exhibit higher levels of aneuploidy in gametes (56–61).

Male rainbow trout exposed to EE2 exhibit high levels of aneuploid sperm but are capable of fertilizing eggs *in vitro*, which leads to the production of high levels of aneuploid embryos. Analysis of NOR staining indicated that chromosome 20 was involved in half of the aneuploidies. Although NOR staining cannot specifically determine hypoploidy/hyperploidy vs. monosomy/trisomy, the natural production of monosomic/trisomic individuals in rainbow trout is an exceedingly rare event when females are properly maintained (62). This fact combined with NOR staining results may indicate that chromosome 20 is somehow specifically susceptible to aneuploidy and/or EE2 exposure during meiosis. As mentioned previously, similar observations have been recorded in human sperm analyses where specific chromosomes are known to have a higher frequency of aneuploidy in normal, fertile males due to decreased meiotic recombination (56–59). This fact may be exacerbated in male rainbow trout because they exhibit the highest degree of recombination suppression in linkage map distance ratios among vertebrates (63), although it remains to be determined if and how estrogens interact and negatively impact recombination sites.

While human autosomal monosomy is considered extremely lethal based on an inability to detect it in clinically recognized pregnancies (64), we observed over two and one-half times the number of hypoploid embryos compared with hyperploid embryos. Several facts may account for this observation in rainbow trout. The first is that salmonid fishes, including the rainbow trout, underwent an autotetraploid event between 25–100 million years ago that duplicated their entire genome (65). Since this event salmonids have undergone Robertsonian rearrangements reducing their chromosome complement (66). Despite these rearrangements, genome duplication can be observed with additional alleles (beyond the normal two) having been identified for many genes (67–69). The presence of such additional alleles in rainbow trout may reduce detrimental effects of embryonic monosomy through some type of compensation, or partial

compensation, which allows embryos to survive longer as a more equal level of transcript formation is maintained (i.e., 4:3 in trout vs. 2:1 in mammal monosomy). The physiological effects of hyperploidy may also be more detrimental than hypoploidy in rainbow trout. Torres and colleagues (70) recently reported in yeast that hyperploidy produced a general phenotype with decreased cell cycle progression and increased environmental sensitivity to protein synthesis and folding, among others. This general phenotype was observed with nearly all of the possible yeast hyperploidy genotypes. Given the reduced number of hyperploidy individuals in our study, an inhibition of growth due to hyperploidy in rainbow trout must be considered.

High levels of aneuploid sperm in EE2 exposed male rainbow trout was strongly correlated with embryonic aneuploidy in this study and was likely the cause of previously observed reductions in embryonic survival (31, 42). This discovery of increased aneuploid sperm formation has broad implications for all sexually mature, or actively spermatogenic, male vertebrates experiencing, or potentially experiencing, environmental estrogen exposure. Despite this advancement in understanding how environmental estrogens affect fertility/development, it remains to be determined what the molecular mechanism is at the level of the developing male germ cell. Because the developing male fish germ cell is sensitive to environmental estrogens, and the fishes are basal in the vertebrate lineage, it is suggestive of a mechanism widely conserved across vertebrate groups.

Materials and Methods

Trout Strain, EE2 Exposure, and Semen Collection. Fish used in these experiments were maintained according to guidelines established by the Institutional Animal Care and Use Committees of Battelle Pacific Northwest National Laboratory, Washington State University and the University of Idaho. Clonal, male Arlee strain rainbow trout (100% homozygous) were used. Fish were propagated at the Washington State University research fish hatchery using androgenesis (i.e., all paternal inheritance) (71). This process results in YY male fish with normal fertility despite lower levels of recombination observed through chromosome markers studies (63, 72). The experimental advantages of clonal rainbow trout are that they are genetically identical and all-male. At 16 months of age (late stage juvenile), 10 individuals were selected from a group previously transported to Marine Research Laboratory (Sequim, WA) and communally held in 370-liter fiberglass tanks with a single-pass flow-through freshwater system with water chemistry/quality parameters previously described (31). Fish were assigned randomly to one of two treatments, five for the control group and five for the chemical exposure group, and placed into 370-liter tanks. Chemical exposures were performed as described in Brown *et al.* (31). Water and stock solution in-flow rates for both treatments were monitored daily and measured by GCMS every 7–14 days. Exposures continued for 50 days after which semen samples were collected from all sexually mature individuals. Fish were anesthetized before sample collection using buffered 0.25 g/l MS-222 (Argent). Individual semen samples were collected by manual expression directly into sterile plastic bags (Whirl-Pak, NASCO) and placed on ice for transport to the University of Idaho. Upon arrival semen was cryopreserved by using standard salmonid sperm cryopreservation techniques with 10 0.5 ml of cryotrays per individual used (73, 74).

Sperm Fluorescent *In Situ* Hybridization (FISH). Three 0.5-ml cryopreserved semen samples from each fish sampled were removed from liquid nitrogen storage and thawed in warm tap water (20°C). The contents of all three cryotrays for each individual were combined in a 15-ml conical tube and fixed with 10 ml of a 3:1 methanol: glacial acetic acid solution. Samples were then centrifuged for 10 min at 2,000 rpm after which time the supernatant was removed and the process repeated two more times. After the final centrifuge cycle, 10 ml of 3:1 methanol: Glacial acetic acid solution was added and samples placed at –20°C or used to prepare FISH slides. Fixed sperm were dropped onto ethanol washed, dd H₂O wetted glass, microscope slides held at a 45° angle and dried over-night before being used in FISH or stored at –20°C. PCR amplification of DNA probes for 18s rDNA (75) and 5s rDNA, which is located on both sex chromosomes in rainbow trout (76) were carried out as reported. The amplified products were labeled with the fluorochrome-labeled dUTPs spectrum green (Vysis, Abbott Laboratories) and spectrum orange (Vysis) for the 18s and 5s rDNA, respectively, using a nick translation kit (Vysis) as described by the manufacturer. Denaturation, probe hybridization

and postwash protocols were carried out as described by the manufacturer (Vysis) with one minor modification. Specifically, no blocking agents (rainbow trout CoT DNA or nonspecies specific DNA) were added with labeled DNA to probe mixtures used for hybridization. After washing, slides were counterstained with DAPI/VECTASHIELD medium (Vector Laboratories) and a glass coverslip applied. Slides were examined using a Leica DMR compound microscope containing a laser light source, appropriate fluorescent filters and attached SPOT camera. A minimum of 500 sperm per individual was analyzed to determine the percent aneuploid sperm per sample.

Fertilization and Embryonic Tissue Preparations. Fertilizations were performed by using standard salmonid *in vitro* fertilization techniques using unfertilized eggs obtained from a single outbred rainbow trout female provided by TroutLodge Inc. and cryopreserved semen. *In vitro* fertilizations were performed by using semen samples from males of both treatment groups (control and EE2 exposed) as described by Cloud *et al.* (77). Nine days after fertilization, developing embryos were prepared for nucleolar organizer region (NOR) and chromosome staining as described in Thorgaard and Disney (78). Embryos were dissected from eggs in 0.9% saline and separated from the yolk before being cultured for 4 h at 19°C in PBS medium containing 25 µg/ml colchicine. After culture, embryos were transferred to a hypotonic solution (1% sodium citrate) for 30 min after which time the hypotonic solution was removed and embryos fixed twice in a freshly prepared 3:1 methanol:glacial acetic acid solution. Embryos were then stored in fixative for a minimum of 12 h at 4°C to ensure complete penetration of fixative. After complete fixation, slides were prepared by using the method described by Kligerman and Bloom (79) with prewarmed slides at 50°C for NOR staining or 45°C for karyotype analysis.

NOR Staining. A single NOR is present in the rainbow trout genome that localizes to chromosome 20 (48, 49). Staining to reveal NORs within the cell nuclei of individuals was performed by using the method proposed by Goodpasture and Bloom (80) and Howell and Black (81) as modified by Gold (82).

This method has been previously used to analyze triploidy in rainbow trout and other salmonid fishes (49). For each individual 50 cells per slide, on two slides (total = 100 cells), were analyzed to determine marker ploidy.

Karyotype Analysis. Slides prepared for chromosome count staining, to determine karyotype, used the protocol described by Thorgaard and Disney (78). After staining slides were evaluated by using Nomarski optics on a Leica DMR compound microscope to identify chromosome spreads suitable for counting. When quality chromosome spreads were identified pictures were taken by using an attached SPOT camera with associated software to obtain digital images for establishment of the karyotype. The karyotype for each embryo analyzed was determined based on the chromosome count from 15 spreads with identical counts from at least 10 required to assign chromosome number. The karyotype for the rainbow trout cross in our study was 62, which is within the range of 58–64 and is consistent with all populations of rainbow trout having 104 chromosome arms (50).

Statistical Analysis. The percent of aneuploid sperm per 500 sperm cells counted was analyzed statistically by using a completely randomized design. The percentage of aneuploid sperm present in each sample was analyzed after performing an arcsine transformation to normalize values. The linear model for this analyses was $y_{ij} = \mu + \alpha_i + e_{ij}$, where α_i represents EE2 treatment effects and e_{ij} the random error. ANOVA was performed to determine whether significant differences were present between treatments using PROC GLM in SAS/STAT (SAS Institute).

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